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FIG. 7A

8con.s = séquence clone pGEM-7/8
pcveco = séquence souche PCV PK/15

SCORES Init1: 2517 Initn: 3774 Opt: 4010
75.8% identity in 1785 bp overlap

1769	1759	1749	1739	1729	1719
8con.s GAATTCTGGCCCTGCTCCCCATCACCCAGGGTGTAGGGAGTGGGCTCCACTGCTGTT					
pcveco GAATTTACCCAGAGACCCATCACCTCTAAATCAAAGAGGTGTTGGTCCACTGTTGTT					
10	20	30	40	50	60
1709	1699	1689	1679	1669	1659
8con.s ATTCTAGATGATAACTTGTAACAAAGGCCACAGGCCCTAACCTATGACCCATATGTAAAC					
pcveco ATCTGGATGCCAACCTTGTAACCCCTCCACCAACTGGCCTATGACCCCTATATTAAC					
70	80	90	100	110	120
1649	1639	1629	1619	1609	1599
8con.s TACTCCTCCGCCATACAATCCCCAACCCCTCTCCTACCACTCCCCTACTTCACACCC					
pcveco TACTCCTCCGCCACACCATAAGGCAGCCCTTACCTACCACTCCAGGTACTCACCCCC					
130	140	150	160	170	180
1589	1579	1569	1559	1549	1539
8con.s AAACCTGTTCTGACTCCACTATTGATTACTTCCAACCAAATAACAAAAGGAATCAGCTT					
pcveco AAACCTGAGCTGGACCAAAACAATTGATTGGTCCACCCAAATAATAAAAGAAACCAGCTG					
190	200	210	220	230	240
1529	1519	1509	1499	1489	1479
8con.s TGGCTGAGGCTACAAACCTCTAGAAATGTGGACCACTGTAGGCCTCGGACTCGCGTTGAA					
pcveco TGGCTCCATTAAATACCCACACCAATGTGAGCACACAGGCCTCGGCTATCGCCTCAA					
250	260	270	280	290	300
1469	1459	1449	1439	1429	1419
8con.s AACAGTATATAC-GACCAGGACTACAATATCCGTGTAAACCATGTATGTACAATTGAGA					
pcveco AA-TGCAGCCACAGCCAAAATTATGTGGTAAGGCTGACTATTATGTACAATTGAGA					
310	320	330	340	350	
1409	1399	1389	1379	1369	1359
8con.s ATTTAATCTTAAAGACCCCCACTTAAACCTAAATGAATAATAAAAACCATTACGAAGT					
pcveco ATTTATCCTAAAGACCCCTC---TAAAT-AAAAATAAAAACCATTACGATGT					
360	370	380	390	400	410
1349	1339	1329	1319	1309	1299
8con.s GAT---AAAAAGACTCAGTAATTATTCATATGGAAATTCAAGGGCATGGGGGGAAAG					
pcveco GATAACAAAAAGACTCAGTAATTATTCATATGGAAAAGGGCACAGGGTGGTCCAC					
420	430	440	450	460	470